

Nonalcoholic fatty liver disease

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Abstract | Nonalcoholic fatty liver disease (NAFLD) is a disorder characterized by excess accumulation of fat in hepatocytes (nonalcoholic fatty liver (NAFL)); in up to 40% of individuals, there are additional findings of portal and lobular inflammation and hepatocyte injury (which characterize nonalcoholic steatohepatitis (NASH)). A subset of patients will develop progressive fibrosis, which can progress to cirrhosis. Hepatocellular carcinoma and cardiovascular complications are life-threatening co-morbidities of both NAFL and NASH. NAFLD is closely associated with insulin resistance; obesity and metabolic syndrome are common underlying factors. As a consequence, the prevalence of NAFLD is estimated to be 10–40% in adults worldwide, and it is the most common liver disease in children and adolescents in developed countries. Mechanistic insights into fat accumulation, subsequent hepatocyte injury, the role of the immune system and fibrosis as well as the role of the gut microbiota are unfolding. Furthermore, genetic and epigenetic factors might explain the considerable interindividual variation in disease phenotype, severity and progression. To date, no effective medical interventions exist that completely reverse the disease other than lifestyle changes, dietary alterations and, possibly, bariatric surgery. However, several strategies that target pathophysiological processes such as an oversupply of fatty acids to the liver, cell injury and inflammation are currently under investigation. Diagnosis of NAFLD can be established by imaging, but detection of the lesions of NASH still depend on the gold-standard but invasive liver biopsy. Several non-invasive strategies are being evaluated to replace or complement biopsies, especially for follow-up monitoring.

Nonalcoholic fatty liver disease (NAFLD) has been accepted as a bona fide disease entity in the late twentieth century. The disorder is characterized by excess accumulation of fat (in the form of triglycerides) in hepatocytes (>5% fat content in the liver; referred to as steatosis), leading to nonalcoholic fatty liver (NAFL). In the presence of additional factors, described below, hepatocyte injury and cell death occur along with lobular and portal inflammation, and the disease is referred to as nonalcoholic steatohepatitis (NASH). With progressive collagen deposition and subsequent vascular remodelling, cirrhosis may result.

Several histological features of NAFLD and NASH are common to alcoholic fatty liver disease (AFLD), including steatosis, inflammation, hepatocyte ballooning (a type of hepatocyte injury), Mallory–Denk bodies (Mallory hyaline) and fibrosis within the lobules (FIG. 1). However, in contrast to the underlying exposure to excess alcohol that initiates and drives AFLD, patients with NAFLD are insulin resistant, usually obese and do not consume excess alcohol¹. Interestingly, whereas paediatric NAFLD shares similar metabolic triggers with NAFLD in adults, different patterns of histological injury may be observed².

Morbidity and mortality in patients with NAFLD can be related to the liver disease itself, although cardiovascular disease is the primary cause of premature death^{3,4}. Cirrhosis and hepatocellular carcinoma (HCC) are the most common liver-related causes of morbidity associated with NAFLD, followed by systemic infection⁵. HCC has been shown to occur in the absence of cirrhosis, which is a concern for clinical care⁶.

Finally, the value of the liver biopsy as a diagnostic tool, although an invasive procedure, continues to be shown by large studies analysing the correlation between long-term outcomes with histopathological evaluation. Advanced fibrosis has been identified as the most important histological feature associated with poor outcomes in two studies^{5,7}; inflammation is associated with the initiation and progression of fibrosis⁸. Both in children and in adults, NASH-related cirrhosis is predicted to become the most common indication for liver transplantation⁹. Several non-invasive diagnostic tools are currently being validated and show promise to replace liver biopsy and identify early fibrosis in the setting of NAFLD.

In this Primer, we describe the worldwide prevalence of NAFLD in adults and children, highlight genomic and epigenomic changes associated with disease and its

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Article number: 15080
[doi:10.1038/nrdp.2015.80](https://doi.org/10.1038/nrdp.2015.80)
Published online
17 December 2015

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progression, as well as the molecular drivers of disease, and elaborate on the advances in diagnosis and the current limitations of management options, quality-of-life (QOL) concerns and outline the avenues for progress.

Epidemiology

Adults

The United States and Europe. NAFLD is currently the most common cause of abnormal liver function tests in western countries. According to population studies using ultrasonography or CT imaging, the prevalence of NAFLD is in the range of 20–50%^{10–13} (TABLE 1). The prevalence of NAFLD varies markedly between ethnic groups. An urban population study in the United States using proton magnetic resonance spectroscopy (¹H-MRS) showed that the prevalence of hepatic steatosis was 45% in Hispanic, 33% in white and 24% in black populations¹⁴. These variations can be explained by differences in lifestyle, the prevalence of metabolic syndrome and genetics, such as a polymorphism of the patatin-like phospholipase domain-containing 3 (*PNPLA3*) gene, which encodes a lipase that mediates triacylglycerol hydrolysis in adipocytes (FIG. 2). NAFLD is strongly associated with metabolic disorders. Not surprisingly, fatty liver has been reported in 40–80% of patients with type 2 diabetes mellitus (T2DM) and 30–90% of patients who are obese^{15,16}.

As imaging can only detect fatty liver but not necroinflammation or fibrosis, the prevalence of NASH and the related liver fibrosis in the population is unclear. Histological analyses suggest that 6–55% of patients with NAFLD have NASH, depending on the patient inclusion criteria and the definition of NASH^{17–19}. Nonetheless, patients undergoing liver biopsy are highly selected and are not representative for a community. In a study of 328 unselected patients attending an outpatient clinic in the southwest of the United States, 46% had NAFLD based on patient history and ultrasonography findings¹². Of those with NAFLD, 134 (out of a total of 156 patients) underwent liver biopsy, of whom 40 were found to have

NASH, suggesting a NASH prevalence of 12% in the adults seeking routine health care. Furthermore, using a score validated against liver histology, a Finnish group estimated the population prevalence of NASH to be around 5%²⁰.

With ongoing necroinflammation and liver injury, patients with NASH can develop progressive liver fibrosis and, eventually, cirrhosis. Advanced fibrosis or cirrhosis due to NASH has a similar mortality than when these features are caused by chronic hepatitis C virus infection and advanced liver disease. Contributing to this risk of death is the development of HCC. The annual incidence of HCC in patients with NASH-related cirrhosis is 1–2%²¹. In the United States, NASH has already become the second-leading aetiology of liver disease requiring liver transplantation in adults because of advanced cirrhosis or HCC²². Importantly, the number of patients with NASH on the transplantation waiting list increased by 170% from 2004 to 2013, the largest absolute and relative increase compared with other aetiologies. Hepatic complications represent the third-most common cause of death in patients with NAFLD or NASH, after cardiovascular events and malignancies⁴.

Asia, Africa and the Pacific Islands. The epidemiology of NAFLD in Asia-Pacific and African countries is shaped by vast differences in lifestyle and economic growth among countries and between urban and rural areas within the same country. For example, in China, the prevalence of NAFLD is higher in coastal and urban areas (up to 30%) than in inland and rural areas (around 1%)²³. In urban areas, the prevalence of NAFLD is around 15–30% in Asia-Pacific countries and 10–20% in Africa.

Compared with western countries, the increase in NAFLD prevalence over time is more marked in Asia and the Pacific areas, probably reflecting rapid lifestyle changes. Studies from Japan and China recorded a two-fold increase in the prevalence of NAFLD over a decade^{23,24}. By contrast, the reported prevalence of NAFLD in the United States was already around 30–40% in 2004 and has not increased substantially thereafter¹⁴. The incidence of NAFLD is difficult to study because ultrasonography cannot reliably quantify liver fat. Nevertheless, according to a recent study from Hong Kong using paired ¹H-MRS, 13.5% of community subjects developed NAFLD in 3–5 years²⁵. Asians and Pacific Islanders develop obesity-related complications at a lower body mass index (BMI) than Europeans and Americans because of the tendency to have central fat accumulation. As a result, approximately 15–20% of Asian patients with NAFLD do not have obesity according to current definitions.

Data on the prevalence of advanced liver disease in the general Asia-Pacific and African populations are scarce. One population study in Hong Kong using transient elastography and serum markers of fibrosis suggests that around 4% of patients with NAFLD in the community have advanced fibrosis or cirrhosis²⁶. In high-risk groups, such as patients with T2DM, advanced fibrosis and cirrhosis can be found in 10–20% of patients.

Children and adolescents

The United States and Europe. NAFLD is the most common liver disease in children and adolescents (TABLE 2). Similarly to adult patients, NAFLD is associated with obesity and metabolic syndrome²⁷ and probably with increased levels of alanine aminotransferase (ALT) in the blood and *PNPLA3* polymorphisms^{28,29} (FIG. 2). A recent analysis of the National Health and Nutrition Examination Survey, including data from >12,000 12–19-year-old Americans, showed that the prevalence of NAFLD (with NAFLD defined as a BMI in the >85th percentile and increased ALT levels of >22.1 U per litre for girls and >25.8 U per litre for boys) more than doubled over the past two decades²⁷. The associated risk factors identified in this study confirm previously identified factors, such as age, increased BMI, male gender and Mexican-American race³⁰. Modifiable factors were

increased consumption of sugar-sweetened beverages and other lifestyle factors (such as a sedentary behaviour)²⁷. Paediatric NAFLD prevalence has been predicted to increase even further³¹.

Ethnicity also has a role in NAFLD risk in children. Given the similar BMI and metabolic syndrome features, black and non-Hispanic children have a lower risk of developing liver disease than Hispanic children (reviewed in REF. 29). Heritability, particularly in first-degree relatives, is documented. Indeed, 59% of siblings and 78% of parents of children with fatty liver will also develop NAFLD³². Age is another striking factor; one post-mortem study in California showed that 17% of teenagers had NAFLD compared with 0.7% of 2–4-year-olds, owing to both a longer period to accumulate steatosis and an increased incidence in adolescents³⁰.

Asia, Africa and the Pacific Islands. Given that overweight and obesity are important risk factors for NAFLD, the prevalence of NAFLD in Korea is greater in patients admitted to obesity clinics (71.9% based on ultrasonography measures and 61.8% based on serum ALT levels) than in general health clinics (8.7% based on ultrasonography measures and 5.9% based on serum ALT levels)³³. In most of Asia, a significant difference is reported between urban and rural populations, with a prevalence of 16–32% in urban areas versus 9% in rural populations³⁴. Obesity-related NAFLD was reported in 64 out of 84 Chinese children (77%) in one study³⁵. In Australia, the prevalence of NAFLD in the paediatric population was estimated to be approximately 10% in the total population and 27.6% among children who are overweight or obese³⁶.

Mechanisms/pathophysiology

Pathophysiology of NAFLD

Fatty liver disease is characterized by hepatic steatosis; there may also be varying degrees of hepatocyte injury, inflammation and, in some cases, fibrosis (FIG. 1).

The evolution of hepatic steatosis in obesity. Several factors contribute to the accumulation of hepatic lipids in the setting of obesity (FIG. 3). First, weight gain is associated with marked expansion of adipose tissue, which leads to dysfunction and eventual death of adipocytes. Dysfunction of adipose tissue results in local inflammation and the upregulation of cytokines that promote insulin resistance. Insulin resistance, in turn, compromises the ability of adipocytes to store fat, resulting in the release of free fatty acids into the circulation, which then become available for uptake by ectopic organs such as the liver³⁷. As a consequence of obesity-related adipose tissue dysfunction, the liver is exposed to high levels of circulating free fatty acids as well as high levels of insulin, which is produced to compensate for systemic insulin resistance. Hepatocytes take up these fatty acids via the transporters fatty acid transport protein 5 (FATP5; also known as bile acyl-CoA synthetase) and CD36, which are also upregulated in obesity^{38,39}. Fatty acid accumulation in hepatocytes prompts the synthesis of triglycerides; during this process, diacylglycerol

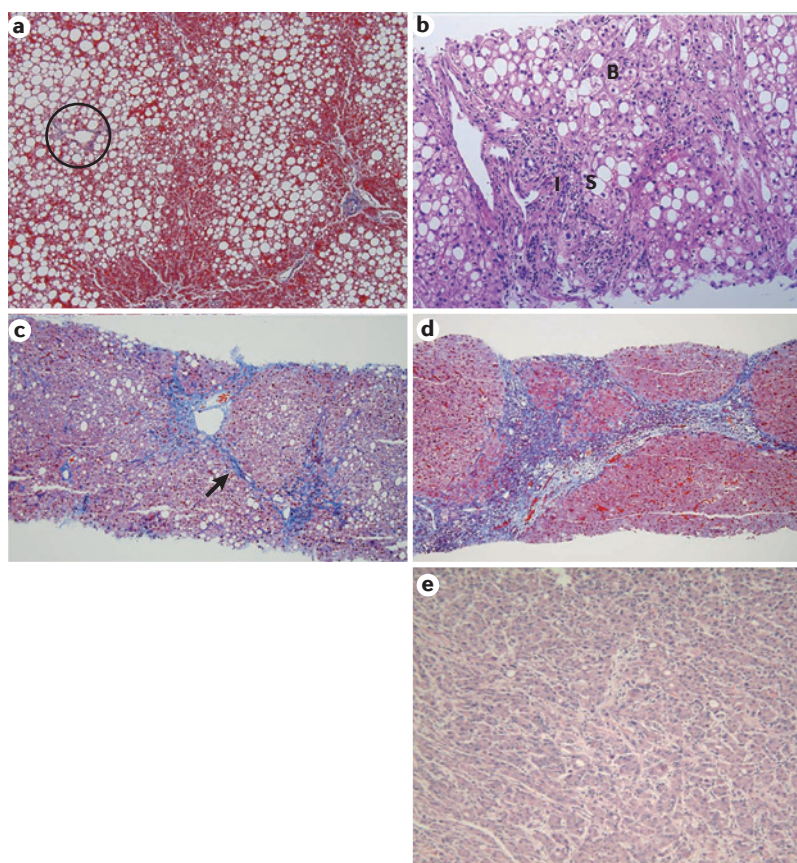


Figure 1 | Histological features of NAFLD. **a** | Marked steatosis without inflammation, hepatocyte injury (ballooning) or fibrosis. Steatosis is concentrated in acinar zone 3, the microcirculatory unit through which blood exits the liver around the terminal hepatic venule (in circle) and shows sparing of the periportal, zone 1 hepatocytes, the microcirculatory unit through which portal and systemic blood enter and mix. This is the adult pattern of nonalcoholic fatty liver disease (NAFLD) (trichrome staining). **b** | Steatohepatitis with marked steatosis (S), ballooning (B), lobular and portal inflammation (I) and extensive bridging fibrosis (haematoxylin and eosin staining). **c** | Fibrosis in the perisinusoidal spaces of zone 3 is detected by trichrome stain; bridging fibrosis (arrow) is noted between two central veins. Hepatocytes with steatosis are seen, but no ballooned hepatocytes are present. **d** | Nonalcoholic steatohepatitis (NASH) with cirrhosis, but no active lesions remain. One would only know this was a case of NASH-related cirrhosis by having had a prior biopsy with the diagnosis of active NASH. **e** | Hepatocellular carcinoma after the development of NASH and cirrhosis.

intermediates accumulate, which impair hepatic insulin signalling by activating protein kinase C ϵ (PKC ϵ)⁴⁰. Hepatocyte insulin resistance fuels gluconeogenesis, promoting hyperglycaemia and prompting even more compensatory insulin production.

The liver itself can contribute to hepatic steatosis by producing lipid from carbohydrate in a process called *de novo* lipogenesis (DNL). The enzymes responsible for DNL are upregulated by insulin and glucose through the action of two transcription factors, sterol regulatory element-binding protein 1 (SREBP1) and carbohydrate-responsive element-binding protein (ChREBP)⁴¹. In the normal liver, DNL is not a main source of hepatic lipid, but in the setting of obesity and hyperinsulinaemia, DNL can contribute as much as 25% of total hepatic lipid stores, and is considered an important factor in the development of NAFLD^{42,43}. Why an insulin-dependent process such as DNL would be upregulated in hepatocytes if uptake of excess circulating fatty acids has rendered them insulin resistant seems paradoxical. One explanation is that hepatocyte insulin resistance manifests downstream from the insulin receptor⁴⁴; another is that DNL can also be induced by insulin-independent pathways, such as endoplasmic reticulum stress-mediated activation of SREBP1 (REF. 45).

Nutrients from the diet also serve as substrates for hepatic triglyceride synthesis and thus participate in the development of hepatic steatosis. Dietary sugars are converted to fatty acids via DNL, and dietary fats are taken up by the liver along with adipose tissue-derived

fatty acids. Roughly 40% of the lipid that accumulates in a fatty liver derives from dietary sugars and fats. Lipolysis of dysfunctional adipose tissue contributes the remaining 60%⁴³.

The gut–liver axis in NAFLD. Interactions between the intestine and the liver are emerging as important determinants of NAFLD. Obesity, which is a key factor contributing to the development of NAFLD, is associated with alterations in the intestinal microbiota that enhance nutrient extraction from the diet⁴⁶. To date, NAFLD has not yet been linked to any specific microbiotic signature, but people with NAFLD have intestinal microbiota that is clearly different from that of healthy individuals⁴⁷. Intestinal dysbiosis can contribute to liver disease by weakening the intestinal barrier and enabling the translocation of bacteria or bacterial products, such as endotoxin or ethanol, into the portal circulation. Dysbiosis can also lead to the luminal degradation of beneficial nutrients, such as choline, which are important for the maintenance of hepatic lipid homeostasis¹¹.

Mechanisms of cell death in a fatty liver. One of the hallmarks that differentiates NASH from NAFL is the presence of hepatocyte injury. Hepatocytes can be damaged by several mechanisms in the setting of NAFLD (FIG. 3). First, the demand for metabolism of excess fatty acids places strain on hepatocyte mitochondria; over time this leads to mitochondrial uncoupling, the production of reactive oxygen species (ROS) and the activation of

Table 1 | **Epidemiology of NAFLD in adults**

Study*	Country or region	Setting	Prevalence of NAFLD (%)	Prevalence of NASH (%)
West				
Autopsy ¹⁶	Canada	Hospital autopsy record	Non-obese: 7 Obese: 29 [†]	Mean: 6 Non-obese: 3 Obese: 19 [†]
Histology ^{17–19}	United States, Europe and Australia	Hospital liver biopsy series	74–96	6–55
Ultrasonography ^{10–13}	United States, Europe and Brazil	General population or hospital clinics	20–70	NR
CT ²³⁴	United States	General population	17	NR
Proton magnetic resonance spectroscopy ¹⁴	United States	General population	34	NR
Asia, Africa and the Pacific Islands				
Histology ^{25,109,235,236}	Korea, Malaysia and Hong Kong	Liver donors and hospital liver biopsy series	Liver donors: 3 Liver biopsy series: 100	56–8
Ultrasonography ^{237–240}	Japan, China, Korea and India	General population and hospital health checkup	9–20	NR
CT ^{241,242}	Saudi Arabia and Egypt	Hospital patients and liver donors	10–18	NR
Proton magnetic resonance spectroscopy ²⁵	Hong Kong	General population	27	NR

NAFLD, nonalcoholic fatty liver disease; NASH, nonalcoholic steatohepatitis; NR, not reported. *When more than one reference is included, the data refer to a summary of several studies. [†]Body mass index of ≥ 30 kg per m².

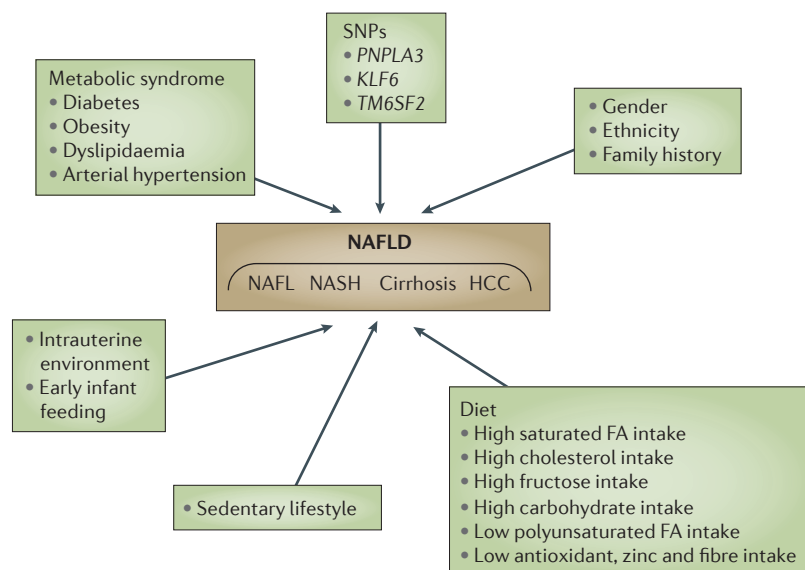


Figure 2 | Factors associated with NAFLD development and progression. Schematic presentations of the factors that influence nonalcoholic fatty liver disease (NAFLD) progression. FA, fatty acid; HCC, hepatocellular carcinoma; *KLF6*, Kruppel-like factor 6; NAFL, nonalcoholic fatty liver; NASH, nonalcoholic steatohepatitis; *PNPLA3*, patatin-like phospholipase domain-containing 3; SNP, single-nucleotide polymorphism; *TM6SF2*, transmembrane 6 superfamily member 2.

Jun N-terminal kinase (JNK)⁴⁸. Once initiated, ROS production and JNK activation continue in a feed-forward loop, which results in mitochondrial damage, impaired ATP production and cell death^{49,50}. Excess fatty acids can also injure hepatocytes by inducing endoplasmic reticulum stress, which leads to mitochondrial dysfunction and cell death through caspase-2-mediated cleavage of BH3-interacting domain death agonist (BID)^{51–53}. However, another pathway via which fatty acids can kill hepatocytes is through the activation of death receptors. Several death receptors (such as FAS, death receptor 5 (DR5; also known as TNFRSF10B and TRAIL-R2) and tumour necrosis factor receptor superfamily member 1A (TNFRSF1A; also known as TNFR1) are upregulated on hepatocytes in the setting of steatosis; the activation of death receptors has been implicated as an important stimulus for hepatocyte apoptosis and necroptosis in NASH^{54–57}.

Inflammation and fibrosis in fatty livers. Activation of the immune system is a key feature of NASH (FIG. 3). The classic effectors of hepatic inflammation in NASH are Kupffer cells and recruited macrophages, but natural killer T cells play an important part in macrophage recruitment⁵⁸, and both natural killer T cells and T cells are emerging as contributors to progressive liver disease^{59–63}. Several compounds can trigger inflammation in fatty livers — including fatty acids, bacterial endotoxin reaching the liver from the intestine and damage-associated molecular patterns (DAMPs) released from dying hepatocytes — and induce inflammation by activating Toll-like receptors (TLRs) and inflammasomes in target immune cells^{64,65}, stimulating the production of an array of cytokines and chemokines^{58,66,67}.

The development of hepatic fibrosis portends a poor outcome in NASH⁷. The central event in hepatic fibrogenesis is the activation of hepatic stellate cells⁶⁸; these cells are susceptible to stimulation by various compounds that are present in a diseased fatty liver. Like inflammatory cells, stellate cells can be activated by DAMPs from dying hepatocytes or other sources that engage TLRs on the cell surface⁶⁹. Other compounds such as free cholesterol can amplify hepatic fibrosis by upregulating the expression of TLRs on stellate cells⁷⁰. Oxidative stress can also activate stellate cells; advanced glycation end products, which are abundant in individuals with diabetes, induce ROS production by stellate cells by inducing the enzyme NADPH oxidase 2 (NOX2; encoded by *CYBB*)⁷¹. One unique consideration in NASH is the role of ballooned hepatocytes in hepatic fibrosis. Ballooned cells produce Sonic Hedgehog⁷², a protein that promotes tissue fibrosis⁷³. Whether Sonic Hedgehog stimulates fibrosis through direct effects on stellate cells or indirect effects on fatty liver injury is currently under study⁷⁴.

Genetics

A substantial proportion of the population are at risk of progressive NAFLD (that is, progression to NASH with advanced fibrosis or cirrhosis) due to obesity and insulin resistance. However, only a minority of people with NAFLD progress to more-advanced disease, including HCC⁷⁵. The reasons for this remain incompletely understood, but NAFLD is best considered a complex disease trait, the development and progression of which are attributed to subtle interindividual variations, including host genetic factors and environmental factors that interact to produce disease phenotype and determine disease progression. Although the presence of NAFLD is principally determined by environmental factors, genetic factors contribute and, crucially, determine how individuals respond to the challenge of caloric excess and consequent metabolic stressors.

Until 2008, studies looking for genetic variants involved in susceptibility to NAFLD had been restricted to case-control allelic association studies limited by the fact that they are reliant on a priori hypotheses for gene selection. Candidate genes are drawn from the limited pool of genes of which biological function is understood and considered relevant to the disease. Only a minority of genes associated with NAFLD through candidate gene analysis have been independently validated in large independent studies or through the use of transmission disequilibrium testing. This short list includes: mitochondrial superoxide dismutase 2 (*SOD2*)⁷⁶, phosphatidylethanolamine *N*-methyltransferase (*PEMT*)⁷⁷, fatty acid desaturase 1 (*FADS1*)⁷⁸, and Kruppel-like factor 6 (*KLF6*)⁷⁹. All of these genes were associated with progressive NAFLD rather than NASH. More recently, several novel genetic associations with NAFLD have been identified through genome-wide association studies (GWAS)^{80–87}. Two of these associations are worthy of particular attention owing to their reproducibility in several GWAS, their association with histologically

Table 2 | Epidemiology of paediatric and adolescent NAFLD

Study (year)	Country	n	Prevalence of NAFLD (%)	Prevalence of NASH (%)
Increased ALT				
Strauss et al. (2000) ²⁴³	United States	2,450	16	NA
Zou et al. (2005) ²⁴⁴	China	113	55.7*	NA
Ultrasonography				
Guzzaloni et al. (2000) ²⁴⁵	Italy	375	38.7	NA
Fu et al. (2006) ²⁴⁶	China	123	80.5	43.9
Chan et al. (2004) ³⁵	China	84	77	24
Histology				
Schwimmer et al. (2003) ²⁴⁷	United States	43	84*	63
Nobili et al. (2006) ²⁴⁸	Italy	84	80*	26.2
Feldstein et al. (2009) ²⁴⁹	United States	66	43	13.7
Autopsy				
Schwimmer et al. (2006) ³⁰	United States	724	13	17.3

ALT, alanine aminotransferase; NA, not available or not stated in the referenced articles; NAFLD, nonalcoholic fatty liver disease; NASH, nonalcoholic steatohepatitis. *In overweight or obese populations.

advanced disease in subsequent association studies and their biological plausibility — at least as genes that are clearly involved in hepatic lipid metabolism.

The first of these genes encodes PNPLA3 (REF. 80). The index single-nucleotide polymorphism (SNP) in *PNPLA3* (rs738409; c.444 C>G; p.I148M) is a non-synonymous cytosine to guanine nucleotide transversion mutation that results in an isoleucine to methionine amino acid change at codon 148. A gene dosage effect for *PNPLA3*^{I148M} carriage is observed with a stepwise increase in ¹H-MRS-quantified hepatic triglyceride content with increasing carriage of the minor allele⁸⁰. The minor allele frequency also correlates with ethnic differences in susceptibility to NAFLD. Multiple histologically based studies have clearly demonstrated that the *PNPLA3*^{I148M} variant is associated with the severity of steatohepatitis and fibrosis⁸⁸ and more recently with the presence of HCC⁸⁹. *PNPLA3* might, therefore, be used as part of an HCC surveillance strategy. The *PNPLA3*^{I148M} SNP has also been associated with a greater response to dietary or lifestyle modification and may therefore be a biomarker for both greater risk of progressive disease and greater benefit from intervention. The precise mechanism through which the *PNPLA3*^{I148M} SNP contributes to disease remains controversial despite intensive study, although alterations in lipid droplet architecture and in retinol metabolism of the hepatic stellate cell have been associated with the *PNPLA3*^{I148M} variant^{90,91}.

The second of these genes encodes transmembrane 6 superfamily member 2 (*TM6SF2*) on chromosome 19, with a non-synonymous SNP (rs58542926; c.449 C>T; p.E167K) associated with ¹H-MRS-quantified hepatic triglyceride content⁸⁶. As with *PNPLA3*, candidate gene studies have shown that the *TM6SF2*^{E167K} variant is also associated with progressive NAFLD⁹². Interestingly, evidence has been provided that carriage of the more

common *TM6SF2*^{E167K} major allele is associated with dyslipidaemia (that is, increased serum LDL cholesterol and triglyceride levels), increased myocardial infarction and cardiovascular disease risk, whereas minor allele carriage is protective⁹³. Functional studies suggest that *TM6SF2* regulates hepatic lipid efflux, with its deletion or mutation resulting in a reduction in lipoprotein secretion (VLDL, triglycerides and apolipoprotein B) leading to increased hepatocellular lipid droplet size and triglyceride accumulation. The *TM6SF2*^{E167K} variant results in lower total cholesterol in humans and expression levels of the variant are lower than the wild-type form. These data indicate that the *TM6SF2*^{E167K} variant confers a 'loss of function' to the protein that may be both qualitative and quantitative⁹³.

Although these studies fail to provide any explanation for the association between the *TM6SF2* variant and progressive NAFLD, they do suggest an interesting paradox — the '*TM6SF2* Catch-22' paradigm⁹⁴. Here, on a background of insulin resistance and metabolic stress, *TM6SF2* genotype acts as a determinant of metabolic syndrome-related end-organ damage and clinical outcome: protecting the liver at the expense of increased risk of atherosclerosis and cardiovascular disease or vice versa. The best evidence in support of this paradigm has recently been provided by a study in patients with NAFLD demonstrating that, although the minor threonine SNP (*TM6SF2*^{E167K}) is associated with advanced disease, carriage of the major cysteine allele (*TM6SF2*^{T167E}) is associated with dyslipidaemia and cardiovascular disease⁹⁵.

Epigenetics

The role of epigenetic factors in the biology of NAFLD is starting to emerge and has enabled integration of information on gene expression with the external and internal metabolic information (FIG. 4). Epigenetic alterations orchestrate the reprogramming of transcriptional machinery of hepatocytes, which occurs during their adaptation to a lipotoxic environment, inflammation and oxidative stress. These epigenetic changes could at least partially explain why the degree of histological damage and/or recovery differs among individuals with an identical genetic background (for example, the presence of the *PNPLA3*^{I148M} risk allele⁹⁶) and in response to similar injury (that is, fat overload).

DNA methylation. DNA methylation in particular is suggested to be involved in the progression of NAFLD towards advanced (fibrotic) clinical stages (FIG. 4). Joined global expression and methylation analysis of human liver specimens demonstrated that advanced disease is associated with global deregulation (mostly hypomethylation) of methylation in multiple CpG sites⁹⁷. Hypomethylation and increased expression of genes involved in tissue repair, regeneration and tumorigenesis were found, as were hypermethylated transcriptionally repressed genes that are involved in basic metabolic functions (such as lipid, urea and amino acid metabolism) or belonged to the cytochrome P450 family⁹⁷. In addition, DNA hypomethylation was associated with

transdifferentiation of stellate cells into myofibroblasts in cell models, a phenomenon commonly associated with fibrosis⁹⁸. Abnormal methylation of the *PNPLA3* promoter was associated with fibrosis severity in a large number of Japanese people⁹⁹. Differences in the methylation status at specific CpG islands within the promoter of anti-fibrogenic (peroxisome proliferator-activated receptor- α (*PPARA*)) and *PPARD*) and pro-fibrogenic (transforming growth factor- β 1 (*TGF β 1*), collagen type I- α 1 (*COL1A1*) and platelet-derived growth factor- α (*PDGFA*)) genes might have a prognostic value in differentiating patients with mild fibrosis from those with advanced fibrosis¹⁰⁰.

Liver-specific epigenetic changes that affect physiological processes and lead to perturbations of normal hepatocyte and mitochondrial function not only affect the progression of NAFLD but also affect NAFLD-related metabolic phenotypes, such as insulin resistance. Targeted analysis of the methylation status of the PPAR γ co-activator 1 α (*PPARGC1A*) — a master transcriptional regulator of glucose and lipid metabolism and mitochondrial biogenesis — showed that methylation levels significantly correlated with plasma insulin levels, β -cell function and the homeostatic model assessment of insulin resistance (HOMA-IR) marker (calculated as (glucose mmol

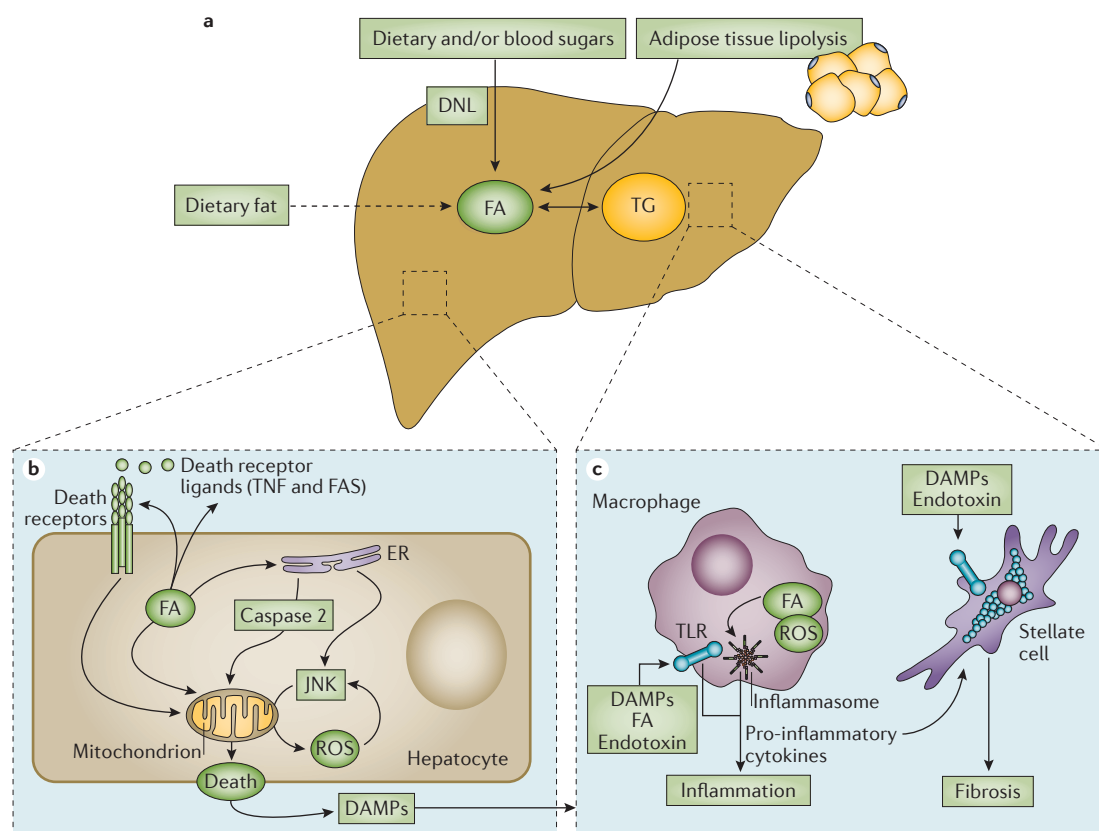


Figure 3 | Mechanisms of steatosis and liver injury in NASH. **a** | Hepatic fatty acids (FAs) come from three sources: diet, *de novo* lipogenesis (DNL) and adipose tissue lipolysis. Under normal circumstances, most FAs come from adipose tissue lipolysis. In people with fatty livers, the absolute amount of FAs from all sources increases, but the proportion coming from DNL increases considerably. Long-chain saturated FAs, which are the products of DNL, are cytotoxic to hepatocytes. The conversion of these FAs to triglycerides (TGs) limits their toxicity. **b** | FAs can kill hepatocytes by upregulating the expression of death receptors and their ligands, culminating in the activation of death receptors ('extrinsic' cell death). FAs can also cause 'intrinsic' cell death by placing stress on the endoplasmic reticulum (ER), leading to caspase-2-mediated cell death, in addition to Jun N-terminal kinase (JNK) activation, reactive oxygen species (ROS) production and mitochondrial dysfunction. In addition, FAs can cause mitochondrial stress directly by creating high demand for β -oxidation. This can also lead to excessive ROS production and mitochondrial dysfunction. Regardless of the pathway involved, FA-induced cell death leads to the release of damage-associated molecular patterns (DAMPs) into the extracellular space. **c** | Hepatic inflammation in fatty livers involves resident and recruited macrophages. These cells express Toll-like receptors (TLRs) and inflammasomes, which respond to DAMPs released from dead hepatocytes as well as other compounds in the local milieu, such as FAs and bacterial endotoxin from the intestine. Activation of these pathways stimulates the production of pro-inflammatory cytokines. Hepatic fibrosis in nonalcoholic steatohepatitis (NASH) is prompted by the activation of hepatic stellate cells to collagen-producing myofibroblasts. These cells express TLRs and, as such, are responsive to DAMPs and endotoxin; they are also stimulated by cytokines released by inflammatory cells. Stellate cells produce collagen, which results in liver fibrosis; they also produce leukocyte chemoattractants (not shown), which can cause a vicious cycle of inflammation and further fibrosis. TNF, tumour necrosis factor.

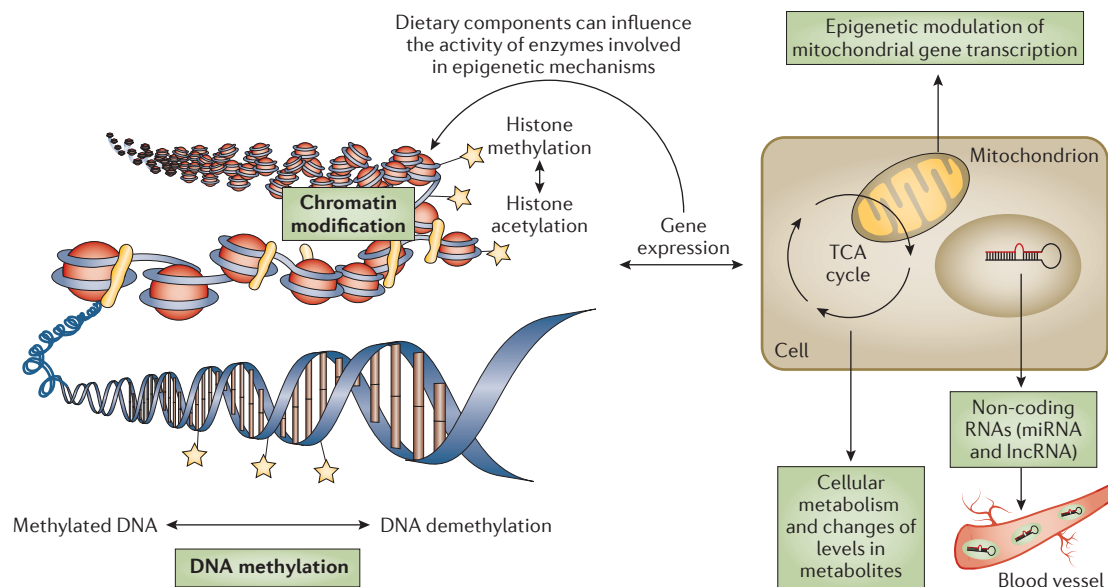


Figure 4 | Overview of factors involved in the modulation of cellular epigenome. The figure depicts overall mechanisms involved in the modulation of cellular epigenome. These epigenetic changes, which are technically regarded as chemical modifications of genomic DNA, include DNA methylation, with methylation leading to gene silencing and demethylation leading to transcriptional activation and post-translational modifications of histones, such as methylation and acetylation. Together, they lead to changes in the chromatin structure that ultimately influence gene expression. Epigenetic alterations can either be transient or permanent; the mentioned epigenetic mechanisms are quite dynamic and can be reversed by therapeutic or lifestyle interventional approaches. Epigenetic changes are shaped by both external and internal influences. For example, dietary components can influence the activity of epigenetic modifications, including histone acetyltransferases, histone deacetylases and histone and DNA methyltransferases. The cellular metabolic environment, including intermediate metabolites and cofactors of the tricarboxylic acid (TCA) cycle, regulate the activity of enzymes involved in histone acetylation pathways and the balance between DNA methylation versus DNA demethylation. Methylation of mitochondrial DNA by DNA methyltransferase 1 can contribute to the pathophysiology of complex diseases such as NAFLD with mitochondrial dysfunction involvement. Finally, non-coding RNAs (including microRNAs (miRNAs) and long non-coding RNAs (lncRNAs)) can regulate transcription, methylation and chromatin structures.

per litre \times insulin mU per ml)/22.5) in humans¹⁰¹. A complex interaction between *PPARGC1A* mRNA levels and liver mitochondrial DNA (mtDNA) copy number was also reported. Interestingly, low mtDNA copy number was also directly associated with insulin resistance and NAFLD¹⁰¹. Methylation of mtDNA might have a role in modulating the liver phenotype beyond genomic DNA¹⁰². Specifically, methylation and transcriptional activation of the mitochondrially encoded enzyme NADH dehydrogenase subunit 6 (encoded by *MT-ND6*), which is involved in oxidative phosphorylation, in the liver was associated with more-advanced liver pathology in humans¹⁰².

A possible explanation for the changes in DNA methylation in NAFLD is its association with obesity, which has been suggested to be associated with reduced DNA demethylation¹⁰³. Furthermore, evidence suggests that the activity of 5-hydroxymethylcytosine (5hmC) might be an additional mechanism that explains liver DNA demethylation pathways in patients with NAFLD¹⁰⁴.

Epigenetic changes can be potentially inherited from one generation to the next. For instance, an experimental study in rodents showed that liver injury induced by the hepatotoxin carbon tetrachloride (CCl_4) led to hypomethylation in the *PPARG* promoter and

associated chromatin modifications in sperm, stem cells and/or mature sperm¹⁰⁵. Although this experimental study suggests that the hepatic wound-healing response might be heritable, it remains unknown whether the same is true in patients with NAFLD. In addition, epigenetic alterations acquired during development may contribute to fetal metabolic programming¹⁰⁶. Maternal protein restriction in mice led not only to fatty liver in the offspring but also to reduced lifespan; further *in vitro* analysis showed that these changes were associated with changes in DNA methylation¹⁰⁷. Likewise, *in utero* overnutrition affects a sirtuin-mediated histone acetylase–deacetylase pathway that has an effect on the hepatic histone code of the progeny, as shown in a non-human primate experimental model of high-fat-diet-induced NAFLD¹⁰⁸.

Non-coding RNAs. A large part (~60%) of the cellular gene expression machinery is regulated by non-coding RNAs — RNA transcripts that do not encode proteins (FIG. 4). MicroRNAs (miRNAs) have been studied extensively in liver disease research, and evidence for their role in NAFLD is consistent. Numerous miRNAs have been implicated in NAFLD severity, liver injury and lipid metabolism, including miR-122, miR-192, miR-375 and miR-34a^{109–111}.

Of particular importance in the pathophysiology of NAFLD is miR-122, the expression of which is highly enriched in the healthy liver. Its network of regulated genes are involved in cholesterol metabolism¹¹², modulation of liver inflammation and progression of hepatocarcinogenesis¹¹³. An *in vitro* study showed that miR-122 regulates the production of collagen in hepatic stellate cells, suggesting a pivotal role in liver fibrogenesis¹¹⁴. Notably, the expression levels of miR-122 are decreased in the liver of patients with NAFLD, whereas levels are increased in the circulating (extracellular) compartment¹¹¹; mouse models of diet-induced steatohepatitis confirmed these findings¹¹⁵. This marked downregulation of liver miR-122 observed in NAFLD seems to be of particular relevance in the pathogenesis of the disease. In fact, silencing of *mir-122* is an early event during hepatocarcinogenesis as a consequence of NASH in humans¹¹⁶. Whether the downregulation of miR-122 in the liver is the cause or the consequence of the progression of NAFLD remains to be investigated, because of the cross-sectional nature of the above-mentioned studies.

Several studies have explored the role of circulating (extracellular) miR-122 levels as a biomarker of NASH — or disease progression — in humans, because miR-122 levels significantly correlate with histological damage (including hepatocyte ballooning and fibrosis) and increased liver enzymes (a marker for liver inflammation or injury)¹¹¹. A two-stage study, exploring a large panel of circulating miRNAs at different stages of NAFLD, showed that circulating levels of miR-122 were increased 7.2-fold in patients with NASH versus healthy controls and 3.1-fold in NASH versus simple steatosis¹¹¹. Although the performance of circulating miR-122 suggests that it could be a reliable biomarker of liver injury in NAFLD, the overall predictive value for liver fibrosis is very similar to currently available non-invasive measurements, including serum cytokeratin 18 levels¹¹¹. Nevertheless, the research agenda is still open; for instance, preliminary data from a rat model of Roux-en-Y gastric bypass showed that circulating miR-122 levels decrease after surgery, suggesting a putative role as a biomarker of response to therapeutic interventions¹¹⁷. Clinical evidence has suggested that bariatric surgery was able to partially reverse epigenetic alterations in the liver associated with NAFLD. Notably, reversing differentially methylated sites in NAFLD correlates with improvement in liver histology¹¹⁸.

Furthermore, miR-122 released from the liver circulate in the bloodstream — like a hormone — and is capable of integrating global metabolic signalling¹¹¹. A functional polymorphism in the 3' untranslated region of a validated miR-122 target gene (that is, rs41318021 of solute carrier family 7 member 1 (*SLC7A1*)) was significantly associated with arterial hypertension, suggesting that miR-122 might potentially modulate the susceptibility to increased blood pressure in people who carry the risk allele of this variant¹¹¹. Moreover, the expression levels of liver miR-122 significantly and negatively correlated with systolic arterial blood pressure¹¹¹. Together, the reduction of miR-122 levels in NAFLD might explain in part the connection between NAFLD and cardiovascular risk.

Finally, although the role of long non-coding RNAs in the pathophysiology of NAFLD remains largely unknown, they are implicated in lipid homeostasis. For instance, a liver-enriched long non-coding RNA in mice regulates lipoprotein lipase activation through a farnesoid X receptor (FXR)-mediated pathway¹¹⁹.

Diagnosis, screening and prevention

Diagnosis of NAFLD

Most patients with NAFLD are asymptomatic or complain about nonspecific symptoms, such as fatigue, sleep disturbances or right upper quadrant discomfort. Hepatomegaly is the most common physical finding. More-advanced liver disease is associated with signs of portal hypertension. Features of polycystic ovary syndrome (associated with hyperandrogenism) should be sought in young women with suspected NAFLD^{120,121}. Obstructive sleep apnoea syndrome and psoriasis can be associated with NAFLD¹²². Apolipoprotein B and lysosomal acid lipase partial deficiency are rare familial causes of NAFLD and should be suspected in the absence of a clear association with components of metabolic syndrome, although lysosomal acid lipase partial deficiency is usually associated with high levels of total and LDL cholesterol^{123,124}.

As no specific marker is currently available, the diagnosis of NAFLD requires the exclusion of all the known other causes of chronic liver disease, haemochromatosis and autoimmune hepatitis and of other causes of steatosis, such as medications (BOX 1), heavy alcohol consumption, viral hepatitis (especially hepatitis C genotype 3), Wilson disease, among others. To discriminate NAFLD

Box 1 | Steatosis and elevated liver tests

Drug use

- Tamoxifen
- Amiodarone
- Glucocorticoids
- Synthetic oestrogens
- Antiviral agents (for example, highly active antiretroviral therapies)
- Methotrexate

Other metabolic or genetic causes

- Hypobetalipoproteinaemia
- Lysosomal acid lipase partial deficiency
- Lipodystrophy
- Weber–Christian disease

Nutrition

- Malnutrition
- Malabsorption
- Total parenteral nutrition
- Rapid weight loss
- Jejunioileal bypass

Others

- Small bowel diverticulosis
- Exposure to petrochemicals
- Exposure to organic solvents

from AFLD, a threshold for daily alcohol intake has been set at <30 g for men and <20 g for women¹²⁵, but a clear differentiation between the two is difficult in those patients who consume a slightly higher amount of alcohol (up to 40 g per day in both men and women).

Another important criterion is that NAFLD is tightly associated with metabolic syndrome and insulin resistance in the liver, muscle and adipose tissues¹²⁶. NAFLD should be suspected in all individuals with one or more components of the metabolic syndrome, defined as the cluster of any three of the following five features: increased waist circumference (ethnicity adjusted), increased fasting glucose or T2DM, hypertriglyceridaemia, low HDL levels (sex adjusted) and increased blood pressure¹²⁷. This association does not characterize NAFLD owing to the genetic variant in *PNPLA3*, but *PNPLA3* mutation and metabolic syndrome may be associated; that is, frequently present in the same person⁸⁰.

Diagnostic workup should include evaluation of family and personal history of components of the metabolic syndrome and assessment of liver tests, including platelet counts, albumin and coagulation, fasting blood glucose, triglycerides and HDL levels. The most common modes of presentation of NAFLD are the detection of unexplained abnormal liver enzymes and/or of bright liver at ultrasonography.

Blood parameters. Liver tests can show mild (twofold to fivefold) increases in the levels of ALT, alkaline phosphatase (ALP) and γ -glutamyltranspeptidase (GGT), but levels are normal in the majority of people with NAFLD (80%)¹²⁸. The ratio of ALT to aspartate aminotransferase (AST) is greater than 1 unless advanced fibrotic NAFLD is present or the patient has covert AFLD. Increased levels of GGT do not discriminate between AFLD and NAFLD, as raised GGT levels are commonly associated with metabolic disease¹²⁹. Ferritin may be increased in up to 60% of patients, but is mainly a marker of subclinical inflammation, given that iron overload is uncommon in NAFLD (4–6%)¹³⁰. However, high ferritin levels (twofold to threefold) have been associated with more-advanced disease¹²⁵. Autoantibodies (antinuclear antibody and α -smooth muscle actin-specific antibody) are present in approximately one-third of patients at low titres (1/80 and 1/160)¹³¹ as compared to absent in healthy individuals.

A test of insulin sensitivity should be considered in all suspected patients. The HOMA-IR is a good surrogate for insulin resistance in individuals who do not have diabetes¹³², although there is no universal agreement on threshold defining what is abnormal. Diagnostic workup should eventually include an oral glucose tolerance test according to standard criteria¹³³ that provides additional information about the glucose tolerance status.

Ultrasound-based techniques. A quantitative ultrasound biomarker for hepatic steatosis is the controlled attenuation parameter (CAP)¹³⁴, which is a quantitative measure of the attenuation (energy loss) of the ultrasound beam as it traverses the hepatic parenchyma. The rationale is that ultrasound energy is attenuated more by fat than by non-fatty tissue; hence, greater attenuation implies

greater fat content. The parameter is measured with the same device used in vibration-controlled transient elastography (VCTE, see below) and is reported in units of dB per metre¹³⁴. An anatomic image is not acquired, therefore, the location where the measurements were made is not recorded. Numerous single-centre studies suggest that CAP can accurately identify patients with hepatic steatosis^{135–137}. However, owing to overlap in CAP values across steatosis grades, CAP has limited accuracy for steatosis grading^{135,136}. In addition, the reproducibility of CAP values in patients with NAFLD has not been studied extensively. Although CAP shows promise, more data are needed before CAP can be considered a valid biomarker of hepatic steatosis.

MRI. The leading quantitative MRI biomarker for hepatic steatosis is proton density fat fraction (PDFF)¹³⁸. This biomarker measures the relative proportion of mobile protons that are attributable to fat and correlates closely with biochemically determined hepatic triglyceride concentration¹³⁸. In numerous single-centre and multi-centre studies, PDFF has shown high accuracy for diagnosing, grading and longitudinally monitoring steatosis in adults and children, including individuals who are morbidly obese^{139–144}. Historically, measurement of PDFF required MRS, which is usually available only in academic centres and generally only assesses portions of the liver. In recent years, advanced MRI techniques have been developed that measure PDFF throughout the entire liver in a single breath-hold, thereby eliminating sampling variability. These techniques are becoming commercially available on the latest generation of magnetic resonance scanners made by the leading manufacturers. Importantly, PDFF measurements made on scanners of different field strength and manufacturer agree closely^{143–145}. With appropriate quality control, these advanced PDFF-estimation techniques can be used in clinical trials (FIG. 5a).

Diagnosis of NASH

The diagnosis of NAFLD can be derived from classic risk factor assessment along with biochemical measures, and ultrasound-based and magnetic resonance-based detection of hepatic steatosis, yet the most relevant challenge to the clinician is the distinction between simple fatty liver and NASH because patients with NASH are at greater risk for developing progressive fibrosis. The presence of normal liver enzymes in the majority (80%) of individuals renders the identification of patients at risk particularly challenging¹²⁸. Importantly, histological severity between patients with and those without abnormal liver aminotransferases does not differ.

The actual diagnosis of NASH requires microscopic evaluation of liver tissue, obtained by liver biopsy (FIG. 1). Histological analysis shows the constellation of steatosis, hepatocyte ballooning and lobular inflammation, typically in acinar zone 3 — the microcirculatory unit through which blood exits the liver¹⁴⁶. Fibrosis better predicts outcomes and mortality⁷. Semi-quantitative grading and staging systems have been developed (BOX 2). These systems are primarily used to assess the severity of lesions. Grading involves the assessment of the activity

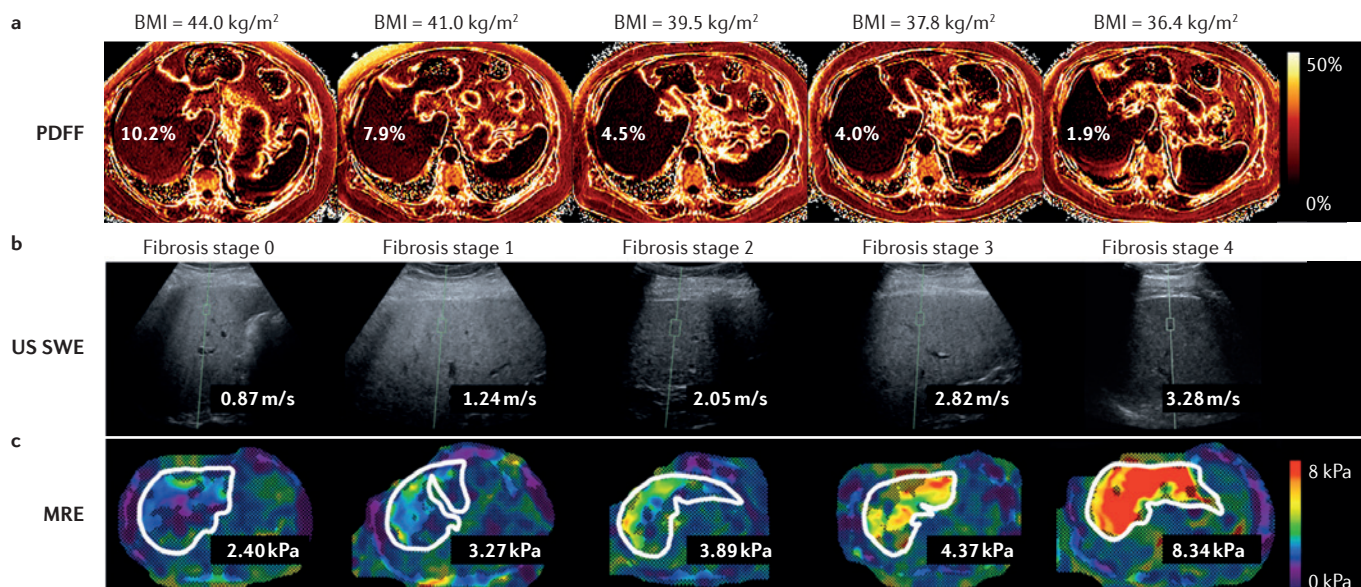


Figure 5 | Diagnosis of liver fibrosis and NASH. Proton density fat fraction (PDFF) maps, obtained by MRI, of an obese woman at multiple time points during a weight loss programme. PDFF is reported as a percentage, with a biological range in humans from 1% to ~50%. Depending on the reference standard, a PDFF cut-off value of 3–6% differentiates non-steatotic from steatotic liver. Notice progressive reduction in PDFF with a reduction in body mass index (BMI; part **a**). Same-day ultrasound-based shear wave elastography (US SWE; part **b**) and magnetic resonance elastography (MRE; part **c**) in five adults with biopsy-proven nonalcoholic fatty liver disease (NAFLD) and fibrosis stages are shown. Notice that patients with NAFLD who also have more fibrosis have greater US SWE-measured shear wave speed and greater MRE-measured stiffness (magnitude of complex shear modulus) than patients who have less fibrosis.

of the disease: steatosis, inflammation and hepatocyte ballooning, whereas staging involves the assessment of fibrosis location and vascular remodelling. Although the grade might be reversible with appropriate intervention, the stage is probably less reversible.

As liver disease progression has been associated with persistence or worsening of metabolic risk factors¹⁴⁷, most predictive indices of disease severity are based on components of the metabolic syndrome along with biochemical and imaging indicators of advanced liver disease. However, the large number of patients with NAFLD and the risk of complications associated with liver biopsy have led to the refinement of non-invasive techniques to predict the histological features of NAFLD and NASH.

In clinical practice, interpretation of non-invasive markers should be performed by hepatologists according to the clinical context and considering all the other clinical and laboratory findings. The combination of two unrelated markers is recommended, as no single test has an advantage over the others in the prediction of the severity of liver disease¹⁴⁸. Among the different strategies, algorithms that combine transient elastography and serum biomarkers are the most attractive and validated¹⁴⁸. However, in case of unexplained discordance of non-invasive tests, a liver biopsy should be performed.

Biomarkers and scores. Screening of steatosis is usually based on ultrasonography in individual patients, but indices or biomarkers are preferred in larger-scale screening studies. The best-validated biomarkers are the Fatty Liver Index, the SteatoTest and the NAFLD liver fat score. Simple steatosis does not increase liver-related mortality,

but these biomarkers can variably predict metabolic and cardiovascular outcomes or mortality¹⁴⁹. Serum markers of fibrosis seem to have a better performance, particularly the NAFLD Fibrosis Score (NFS), Fibrosis-4 (FIB-4), BARD and commercially available panels, such as FibroTest, FibroMeter and the Enhanced Liver Fibrosis (ELF) test¹⁵⁰ (TABLE 3). All of these tools have acceptable diagnostic accuracy, but only NFS and FIB-4 have been extensively validated¹⁹. All of these tests perform best at excluding severe fibrosis–cirrhosis, with negative predictive values of >90%, but are typically less accurate in the determination of less-severe fibrosis¹⁵⁰. NFS, FIB-4, ELF and FibroTest can also predict overall, cardiovascular and liver-related mortality¹⁴⁹.

Several non-invasive procedures have been tested for the non-invasive prediction of NASH. Plasma levels of cytokeratin 18 fragments (a marker of apoptosis) had a promising performance, but recent data have highlighted the limited accuracy¹⁵¹ and to date no recommendation can be issued for the non-invasive diagnosis of NASH.

Ultrasonography. Conventional ultrasonography, in which standard B-mode (grey scale) ultrasonographic images are acquired and interpreted qualitatively, is frequently used to screen for hepatic steatosis, but conventional ultrasonography is insensitive to mild steatosis, especially in individuals who are obese, and is limited by reader variability^{152,153}.

Elastography. Imaging techniques that are collectively known as elastography have been developed to indirectly measure tissue stiffness non-invasively.

Box 2 | Grading and staging histological features of NAFLD and NASH

NAFLD Activity Score (NAS; 0–8)*

- Steatosis (0: <5%; 1: 5–33%; 2: 33–66%; and 3: >66%)
- Lobular inflammation, foci per 20× magnification (0: not present; 1: <2; 2: 2–4; and 3: >4)
- Ballooning (0: not present; 1: few; and 2: prominent ballooning)

NAFLD Fibrosis Score (0–4)*

- 1a: delicate zone 3 psf[†]
- 1b: dense zone 3 psf[†]
- 1c: portal only
- 2: zone 3 plus portal or periportal
- 3: bridging (c–c, c–p and p–p)
- 4: cirrhosis

Fatty Liver Algorithm[§]

- Steatosis (0–3)
- Activity (ballooning and lobular inflammation)
- Ballooning (0–2)
- Lobular inflammation (0–2)
- Fibrosis (similar to the fibrosis score of the NASH CRN above)

c–c, central-to-central; c–p, central-to-portal; CRN, Clinical Research Network; NAFLD, nonalcoholic fatty liver disease; NASH, nonalcoholic steatohepatitis; p–p, portal-to-portal; psf, perisinusoidal fibrosis. *National Institute of Diabetes and Digestive and Kidney Diseases (NIDDK) NASH CRN: Pathology Committee NAFLD Scoring System for Clinical Trials²²⁷. Although the NAFLD Activity Score is known to closely correlate with a diagnosis of NASH, it was created for clinical trials to assess changes in the components and has been shown to be associated with liver tests (alanine aminotransferase and aspartate aminotransferase), whereas the pathologist's diagnosis of NASH associates with features of insulin resistance. The pathologist is encouraged to make a separate, pattern-based diagnosis in addition to scoring the lesions. [†]Delicate psf requires trichrome staining, whereas dense psf can be visualized on a haematoxylin and eosin stain first. [§]A primary difference between the European algorithm and the NASH CRN is that the former score was derived to establish a diagnosis of NASH. Steatosis of >1, activity of >2 (with both ballooning and lobular inflammation of >1) equals NASH²²⁸.

The measurement involves the assessment of the propagation of shear waves within the liver, from which the hepatic stiffness is inferred. Depending on the device, results can be reported as shear wave speed in metres per second or as one of several elastic moduli (for example, Young modulus or complex shear modulus) in kPa¹⁵⁴. These various elastography-derived parameters have emerged as the leading imaging biomarkers of hepatic fibrosis, but like serum markers, they perform best only at detecting advanced (stage 3 or stage 4) fibrosis.

Ultrasound-based elastography. The first ultrasound-based method to measure liver stiffness was VCTE¹⁵⁵. This technology uses a non-imaging device to track shear waves that are transmitted transiently into the liver by an external vibration source; an anatomical image is not acquired and the location from which the measurements are made is not recorded.

More recently, elastography capabilities have been added to clinical ultrasound scanners¹⁵⁴. Elastography-enabled scanners use ultrasound to track shear waves that are generated transiently *in situ* within the liver by an ultrasound push pulse (also known as an acoustic radiation force impulse); measurement locations are recorded on an anatomical image^{154,156,157} (FIG. 5b). As each manufacturer uses proprietary hardware and software, measurements made with different devices and even with different

transducers (probes) on the same device do not agree^{157,158}. Thus, any given patient should be monitored serially with a single device and using the same probe. VCTE and elastography-enabled scanners have been shown to provide high accuracy for the diagnosis of advanced fibrosis in patients with viral hepatitis infection^{159,160}. Single-centre studies suggest these devices also accurately diagnose advanced fibrosis in patients with NAFLD¹⁶⁰, although multi-centre validation studies have not yet been performed. None of these devices can diagnose early fibrosis accurately. The technical success rate, reliability, robustness and precision of these devices for measuring stiffness parameters in NAFLD has not been extensively studied, and their performance in detecting changes in fibrosis in the context of treatment trials is unknown¹⁵⁷. Until such data become available, caution is advised for use of these devices in NAFLD clinical trials.

Magnetic resonance-based elastography. Magnetic resonance elastography uses a specialized MRI sequence to visualize shear waves that are delivered continuously into the liver by an external acoustic source¹⁶¹. The wave images are processed by a so-called inversion algorithm to generate 'elastograms' (REF. 162) (FIG. 5c). These computer-generated images depict the spatial distribution of a stiffness parameter known as the magnitude of the complex shear modulus (usually referred to as 'shear stiffness' in the medical literature). The hepatic stiffness is recorded from representative portions of the liver. Magnetic resonance elastography is now commercially available on commercial MRI scanners, with the same hardware to generate the shear waves and the same inversion algorithm to process the wave images. Small single-centre studies in healthy volunteers suggest that measurements made with different scanners agree¹⁶³. Large studies in patients with NAFLD are needed to confirm cross-platform reproducibility.

In single-centre studies, magnetic resonance elastography has been shown to diagnose advanced fibrosis in adult NAFLD with high accuracy^{164–166}. The accuracy of magnetic resonance elastography is probably higher than that of ultrasound-based elastography¹⁶⁷. As with ultrasound-based techniques, accuracy for the detection of early fibrosis is modest. The precision of magnetic resonance elastography in the NAFLD population has not been studied extensively. Multi-centre validation studies are needed before magnetic resonance elastography can replace biopsy in NAFLD clinical trials.

Follow-up monitoring

Patient monitoring should include routine biochemistry, non-invasive monitoring of fibrosis and the assessment of co-morbidities. Given that the risk of developing T2DM and cardiovascular complications is increased in patients with NAFLD¹²², oral glucose tolerance tests should be performed whenever glucose metabolism is abnormal (that is, a fasting glucose of 100–126 mg per dl; glycated haemoglobin (HbA1c) of 5.7–6.4%)¹³³ and carotid doppler ultrasonography¹²²; referral to other specialists might be required. The timing of follow-up testing is still debated, and a definite evaluation for risks

of progression and costs associated with investigations has not yet been performed. In patients with simple fatty liver, a reasonable approach is a follow up in primary care setting, unless worsening of metabolic risk factors occurs. In patients with NASH and/or fibrosis, a yearly monitoring policy is advisable. If indicated, on a case-by-case basis a repeat biopsy could be performed after 5–7 years.

Management

Current management of NAFLD

The first step in the management of NASH in most patients is the implementation of lifestyle modifications that focus on healthy eating habits and regular exercise¹⁶⁸. Without biopsy confirmation of the presence of NASH, lifestyle modification is still appropriate to reduce cardiovascular risks in patients with NAFLD and also to treat NASH if it is present. Weight loss for patients who are overweight and obese is recommended based on the collective observations from multiple studies showing that histological improvement can occur when patients lose 5–10% of their body weight^{169,170}. Unfortunately, meaningful lifestyle modification is difficult to achieve and difficult to sustain for most people^{171,172} and,

therefore, alternatives are needed to treat NASH and to prevent the associated complications of the metabolic syndrome. At this time, no pharmaceutical treatments are approved for NASH, although many agents are in various stages of evaluation in clinical trials. Bariatric surgery is a good option for select patients who are obese and have NASH as it typically leads to improvement in NASH^{173,174}.

Pathogenesis-based targets of therapy

As reviewed above, one perspective on the pathogenesis of NASH is that lipotoxic lipid species, which are derived from an excess supply of fatty acids in hepatocytes, mediate cellular injury and the resulting inflammatory and fibrotic response that results in the histological phenotype of NASH^{175–177}. This paradigm enables us to predict therapeutic interventions that might prevent or treat NASH by targeting the oversupply of fatty acids to the liver, the generation of lipotoxic intermediates and the resulting dysfunctional cellular responses (FIG. 6). Given that the best treatments of disease are those that target underlying causes at a fundamental level, an ideal approach would be to pharmacologically manipulate the pathways of energy metabolism that are responsible for the heterogeneity in human energy efficiency.

Table 3 | Scores predicting NAFLD and NAFLD severity (fibrosis)

Score	Components	Cut-off value	AUROC	Sensitivity (%)	Specificity (%)	PPV	NPV	Notes
NAFLD								
Fatty Liver Index ²⁵⁰	BMI, WC, TG and GGT	>60	0.85	61	86	NA	NA	Less accurate in Asian population Predicts metabolic and CV outcomes, and hepatic and CV mortality
NAFLD liver fat score ²⁵¹	MetS, T2DM, AST and ALT	−0.640	0.86	86	71	NA	NA	Prediction not improved by PNPLA3
SteatoTest® (REF. 252)	GGT, ALT, BG, TG and CHOL	>0.69	0.80	38	81	71	52	Predicts overall mortality
Fibrosis								
NAFLD Fibrosis Score ²⁵³	Age, BG, BMI, platelets, albumin, and AST or ALT	≥0.676	0.84	43	96	82	80	Predicts liver-related events, incident diabetes, all-cause and CV mortality
Fibro-Test® (REF. 254)	GGT, BIL, haptoglobin, apoAI and α2-macroglobulin	>0.30 >0.70	0.81	92 25	71 97	33 60	92 89	Predicts overall mortality, progression to advanced fibrosis; associated with CV mortality
FIB-4 index ²⁵⁵	Age, ALT, AST and platelets	≥2.67	0.80	33	98	80	83	Predicts all-cause and CV mortality and liver-related events
BARD ²³²	BMI, AST or ALT and T2DM	2–4	0.81	NA	NA	43	96	Predicts liver-related events
Hepascore ²⁵⁶	Age, gender, α2-macroglobulin, HA, BIL and GGT	>0.37	0.81	75	84	57	92	Initially developed in hepatitis C virus
Enhanced Liver Fibrosis ²⁵⁷	HA, TIMP1 and PIIINP	>0.3576	0.90	80	90	71	94	Predicts overall and CV mortality
AST/platelet ratio index ¹⁴⁸	AST levels and platelet counts	≥0.918	0.87	66	91	73	87	Predicts liver-related events

Generally, the performance of a non-invasive diagnostic method is evaluated by the area under the receiver operator characteristic curve (AUROC), taking liver biopsy as the reference standard, in which a value of 1 reflects a 'perfect' test with 100% sensitivity and specificity and a value of 0.5 reflects a test as good as chance. ALT, alanine aminotransferase; apoAI, apoprotein AI; AST, aspartate aminotransferase; BG, fasting blood glucose; BIL, total bilirubin; BMI, body mass index; CHOL, fasting cholesterol; CV, cardiovascular; GGT, γ-glutamyl-transpeptidase; HA, hyaluronic acid; MetS, metabolic syndrome; NA, not available; NAFLD, nonalcoholic fatty liver disease; NPV, negative predictive value; PIIINP, amino-terminal propeptide of type III collagen; PNPLA3, patatin-like phospholipase domain-containing protein 3; PPV, positive predictive value; T2DM, type 2 diabetes mellitus; TG, fasting triglycerides; TIMP1, tissue inhibitor of metalloproteinase 1; WC, waist circumference.

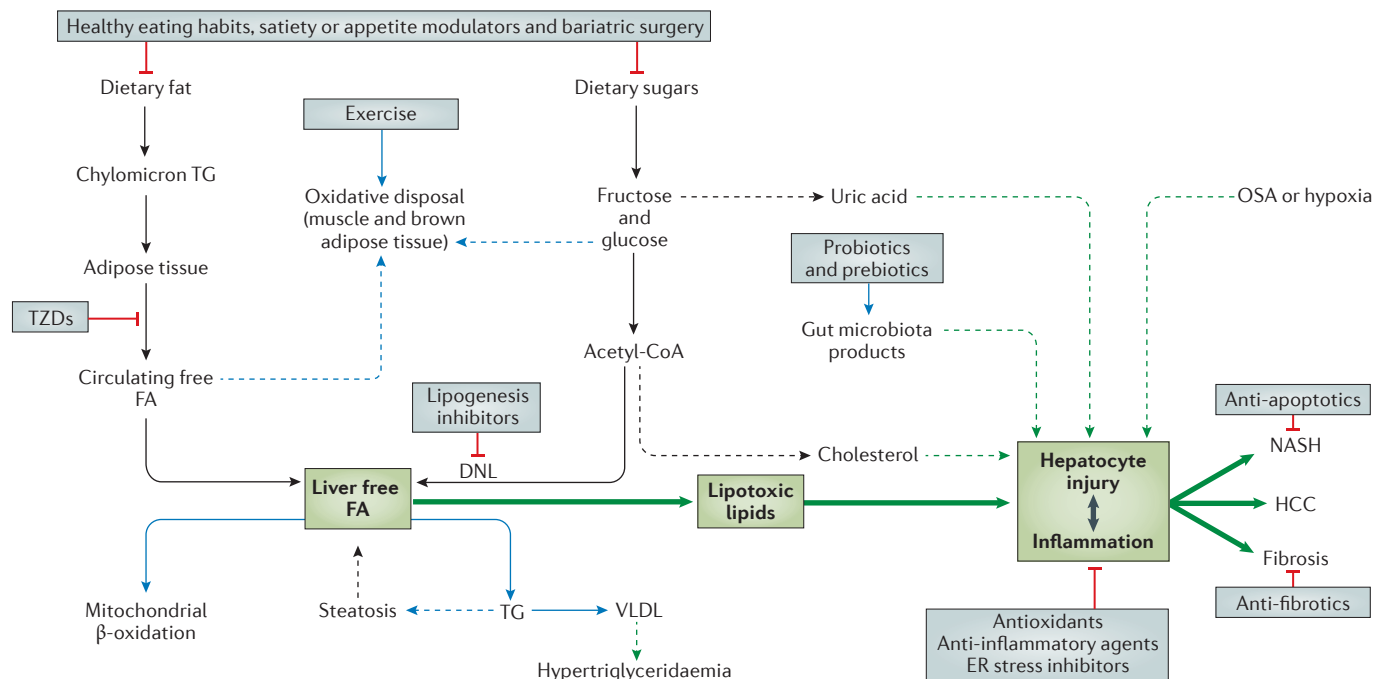


Figure 6 | The lipotoxicity model of NASH and targets for therapy. Central to the understanding of the pathogenesis of nonalcoholic steatohepatitis (NASH) are free fatty acids (FAs), their sources in hepatocytes and their fates. Delivery from adipose tissue and *de novo* lipogenesis (DNL) are the main sources of free FAs in hepatocytes. Mitochondrial β -oxidation and secretion into the blood as triglycerides (TGs) in VLDL are the major fates. When the supply is excessive or the disposal is impaired (or most likely a combination of both), the generation of lipotoxic lipids can occur. This leads to hepatocyte injury, inflammation and the phenotype of NASH. These processes can be a stimulus for fibrogenesis and carcinogenesis. This paradigm predicts therapeutic targets (grey boxes), many of which are being explored in clinical trials. Black arrows indicate normal substrate flow; green arrows indicate flow or processes that contribute to NASH and its consequences; blue arrows indicate flow or processes that can prevent NASH and its consequences; and dashed arrows indicate pathways that can be important in substrate flow under specific circumstances. ER, endoplasmic reticulum; HCC, hepatocellular carcinoma; OSA, obstructive sleep apnoea; TZD, thiazolidinedione.

If the genomic or epigenetic changes responsible for these differences can be identified and exploited, then treatments could be effective for NASH as well as the underlying insulin resistance, hypertension, hyperlipidaemia and other manifestations of the metabolic syndrome that lead to disability and early death¹⁷⁸. Animal studies have provided some insights into these differences, but how these will be translated into treatment of human disease is yet to be shown. Examples include the divergence of rat strains into those having high and low exercise tolerance, with features of the metabolic syndrome developing in the latter¹⁷⁹, the role of β -adrenergic stimulation of highly metabolic brown adipose tissue¹⁸⁰, the role of serotonin in modulating metabolism¹⁸¹ and the role of peripheral cannabinoid signalling in modulating energy efficiency¹⁸². At this time, insights into the regulation of energy metabolism are mostly at early preclinical stages, and clinical trials have generally focused on manipulating more downstream processes, such as lipogenesis, lipolysis, inflammation, oxidative stress and fibrogenesis.

Preventing an oversupply of fatty acids in the liver
Inhibition of DNL. One approach to reducing lipotoxic liver injury is to reduce the endogenous production of fatty acids in the liver by inhibiting DNL. The transcription factor SREBP1C is a major regulator of DNL; thus,

treatments that decrease SREBP1C may be beneficial. This is one potential mechanism by which the bile acid derivative obeticholic acid may work. As an activating ligand for the nuclear receptor FXR, obeticholic acid increases the expression of tyrosine-protein phosphatase non-receptor type 6 (PTPN6; also known as SHP1), which decreases the expression of SREBP1C¹⁸³. A Phase II clinical trial of obeticholic acid has demonstrated histological improvement in NASH after 72 weeks (45% versus 21% in placebo-treated patients; $P=0.0002$)¹⁸⁴. However, treatment was associated with significant pruritus in 23% in the treatment group compared with 6% in the placebo group, and with an increase in total and LDL cholesterol and a decrease in HDL cholesterol in treated patients, the importance of which will be further assessed in another clinical trial.

Another approach to inhibiting DNL includes using inhibitors of key enzymes in the pathway of fatty acid synthesis, such as acetyl-CoA carboxylase, and such agents are being evaluated in clinical trials. Augmenting the incretin axis with glucagon-like peptide 1 (GLP-1) analogues such as exenatide or impairing the breakdown of endogenous GLP-1 with the dipeptidyl peptidase 4 inhibitors (gliptins) may also decrease hepatic lipogenesis by decreasing substrate availability; this approach is also being examined in clinical trials.

Reducing the burden of fatty acids delivered to the liver. Lipolysis of stored triglyceride in adipose tissue is the main source of fatty acids delivered via the circulation to the liver. Adipose tissue lipolysis is normally suppressed by insulin, but when adipose tissue is insulin resistant, it continues to release fatty acids into the blood at inappropriate times and thus contributes to the development of NASH¹⁸⁵. The PPAR γ ligands thiazolidinediones (TZDs or 'glitazones') have been shown to improve adipocyte insulin responsiveness and thus decrease inappropriate lipolysis. One TZD, pioglitazone, has been shown to improve adipose insulin sensitivity and improve NASH^{186,187}. Unfortunately, this benefit comes at the expense of increased body weight in some patients because of adipocyte triglyceride retention. This and possibly other adverse effects such as osteoporosis and bladder cancer have dampened the enthusiasm for using pioglitazone for NASH and it has not gained widespread use.

Metformin is another insulin sensitizing agent but its effect is primarily in the liver. There is little evidence that hepatic insulin resistance has a role in causing NASH. A large randomized clinical trial (RCT) of metformin in children did not show any benefit on ALT levels or histology but fortunately neither did it show worsening¹⁸⁸. Nonetheless, metformin may have an adjunctive role in the treatment of NASH because it is often associated with weight loss, and recent data indicate that it is associated with a reduced risk of several malignancies¹⁸⁹.

Fatty acids and the carbohydrates used to make fatty acids can also be diverted to oxidative pathways and this may explain the benefit of regular exercise in improving the metabolic syndrome. Oxidative pathways can also be stimulated pharmacologically. The PPAR α and PPAR δ ligand GFT505 was recently evaluated in a clinical trial because it improves insulin sensitivity, which may be explained by the effect of PPAR α in promoting oxidative metabolism in the liver and PPAR δ in promoting oxidative metabolism in muscle¹⁹⁰.

Reducing injury

The mechanisms of lipotoxic injury remain uncertain, but oxidative stress has been proposed to have a role¹⁹¹. Oxidative stress with associated lipid peroxidation clearly occurs in NASH¹⁹², but whether it is causative or an epiphenomenon remains unknown. Two large RCTs of natural vitamin E (d- α -tocopherol) at a dose of 800 IU daily have been conducted: one in adults (PIVENS) and one in children (TONIC). The PIVENS trial demonstrated improvement in liver histology in 43% compared with a placebo response of 19% ($P = 0.001$)¹⁸⁷, and similarly the TONIC trial showed resolution of NASH in 58% compared with 28% in placebo ($P = 0.006$)¹⁸⁸. Neither trial showed evidence that indices of oxidative stress improved, so it remains somewhat unclear how vitamin E improves NASH. Both the PIVENS and the TONIC trials were conducted in patients without diabetes or cirrhosis, therefore, the benefits of vitamin E in patients with these commonly associated conditions remain unknown. Concern has been raised about the cardiovascular safety of high-dose vitamin E and,

consequently, patients and doses should be carefully selected. The ideal patient may be someone with aggressive NASH and no diabetes or major risks for cardiovascular disease, but such patients comprise a relative minority. It should be noted that weight loss has an additive benefit in patients treated with vitamin E, so the use of vitamin E should be an adjunct to weight loss and not an alternative¹⁹³. Fish oil (poly-omega-3 fatty acids) may also exert beneficial effects on lipid peroxidation, but a large trial of ethyl-eicosapentaenoic acid in patients with NASH did not show any benefit¹⁹⁴.

Endoplasmic reticulum stress is another process known to occur in NASH that may be central to the pathogenesis of lipotoxic hepatocyte injury. Studies of agents that diminish the endoplasmic reticulum stress response have provided provocative data in animals but these have yet to be introduced into RCTs¹⁹⁵.

The contribution of hypercholesterolaemia to the development of steatohepatitis and liver fibrosis has not been fully established, but several studies suggest that it may have a role in some patients. A morphometric study of liver biopsies demonstrated cholesterol crystals in lipid droplets in patients with steatohepatitis but not steatosis¹⁹⁶. Statin use is safe in patients with liver disease¹⁹⁷, and a *Cochrane* review concluded that statins can improve steatosis and reduce the levels of aminotransferase, but treatment trials with histological endpoints are lacking¹⁹⁸. More recently, in a large European observational cohort with NAFLD, statin use was associated with less steatosis, NASH and fibrosis in a dose-dependent manner; interestingly, the presence of the *PNPLA3*^{148M} polymorphism prevented the protective effect of statins¹⁹⁹. In another recent study, statin use in patients with NAFLD was associated with a reduction in liver-related deaths, liver transplantations or other liver outcomes⁷.

Inhibition of inflammation

The role of inflammation in contributing to hepatocyte injury versus a consequence of injury is unknown but should be clarified by the results of clinical trials examining various anti-inflammatory approaches. One clinical trial of an anti-inflammatory phosphodiesterase type 4 inhibitor was negative despite the demonstration of reduced circulating tumour necrosis factor levels in patients given the active agent²⁰⁰. However, like all contributors to the pathogenesis of NASH, inflammation may have an important role in only a subgroup of patients. Post-hoc analysis of RCT results to identify and then predict responders to a specific therapy will be essential for all agents.

Inhibition of apoptosis

Caspase inhibitors have been developed and are being explored in RCTs for numerous diseases including NASH based on the recognized role of hepatocyte apoptosis in the pathogenesis of NASH²⁰¹. Although this might be a rational approach in acute processes, theoretical concerns about the safety of long-term inhibition of apoptosis persist with this treatment approach in NASH.

Inhibition of fibrosis

In the absence of effective therapy for NASH, being able to prevent the progression of fibrosis to cirrhosis would be advantageous. Even when effective NASH therapies are identified, there may be a role for short-term use of effective anti-fibrotic agents to accelerate the reversal of fibrosis. No anti-fibrotic agents for liver disease are available but several are being explored in RCTs, including an anti-lysyl oxidase homologue 2 (LOXL2) antibody and a galectin-binding molecule that have shown early promise.

Quality of life

QOL in patients with NAFLD is understudied, although its importance is substantial and broadens our understanding of the overall burden of the disease (BOX 3). Several studies have reported QOL scores in the context of NAFLD^{202–204}. Overall findings have been similar and suggest that NAFLD is associated with a reduced overall but predominately physical health-related QOL (HRQOL). Some data suggest that the impairment in QOL reported in patients with NAFLD may be beyond that reported in other aetiologies of liver disease^{202,203}. In a sample of 771 patients with biopsy-confirmed NAFLD from the NASH Clinical Research Network database, QOL as measured by the Short-Form 36 Health Survey questionnaire was compared to a United States reference population with or without chronic disease²⁰⁵. In this study, impaired QOL was most evident in physical health, with mental health affected to a lesser degree.

Most patients with NAFLD are overweight or obese, and even in those who are not, lifestyle changes that incorporate dietary modification and exercise are the cornerstone of therapy. Obesity has both health and psychosocial ramifications and can have a profound effect on an individual patient's QOL. In a cross-sectional study, obesity was not associated with lower QOL²⁰⁵; however, the effects of imposing lifestyle change were not measured. One study evaluated the effect of weight loss on HRQOL in patients with NAFLD and found that, compared with those who did not lose weight, those that did had improved HRQOL scores over baseline values, comparable to the general population²⁰⁴. A prospective

study of intensive lifestyle intervention showed that patients who lost >7% body weight benefited not only from improvement or resolution in steatosis, necroinflammation and cellular ballooning (NASH resolution) but also improvement or resolution in fibrosis¹⁷⁰. Importantly, no patient who lost >7% body weight had progression of fibrosis. In those that lost >10% body weight, fibrosis improved in 81%, although overall numbers were small as was baseline extent of fibrosis. Weight loss and dietary change are hard to achieve and even harder to maintain, particularly in the case of morbid obesity. Not only is behavioural change difficult in itself but also barriers — physical, social and economic (among others) — can pose a challenge to accomplishing them effectively.

Bariatric surgery for NASH

Compared with patients who are obese and are matched for co-morbidities, those that undergo bariatric surgery have decreased long-term mortality^{206–208}. The largest effect on mortality comes from a reduction in myocardial infarction and malignancy — the two most common causes of death in patients with NAFLD. Bariatric surgery can be an effective and appropriate treatment strategy for a subset of patients with NAFLD, including those with advanced fibrosis and even carefully selected patients with compensated cirrhosis who meet criteria for bariatric surgery^{4,209}. While it is important to note that bariatric surgery has not been studied prospectively specifically as a treatment for NASH, the best-available data demonstrate that bariatric surgery in patients with NASH is safe and improves NASH-related liver disease, including fibrosis¹⁷⁴. The liver-related benefits derived from bariatric surgery such as Roux-en-Y gastric bypass or sleeve gastrectomy extend beyond weight loss. Both procedures increase the levels of GLP-1, which decreases appetite, slows gastric emptying and improves insulin sensitivity. Furthermore, GLP-1 affects the modulation of bile acid signalling, specifically through FXR, which can alter the gut microbiota and can have other potential benefits^{210–212}.

However, practitioners and patients are reluctant to pursue bariatric surgery, even when patients are appropriate candidates. Although there are known complications that vary according to the procedure performed, the overall risk is low in properly selected patients²¹³. Although carefully selected patients with established compensated (that is, stable without profound abnormalities in liver function) cirrhosis can safely undergo bariatric surgery, cirrhosis is associated with a higher risk of postoperative complications, including hepatic decompensation (that is, the development of ascites, hepatic encephalopathy or variceal bleeding)²⁰⁷. Therefore, careful patient selection cannot be overemphasized. Success of the intervention requires that patients remain committed to necessary lifestyle changes postoperatively to avoid regaining lost weight. More importantly, compliance with recommendations and follow-up are imperative to avoid metabolic and nutritional problems that are related to malabsorption or other consequences of bariatric surgery²¹⁴.

Box 3 | Affected quality-of-life aspects in NAFLD

Physical health scores

- People with nonalcoholic steatohepatitis (NASH) have poorer scores than patients with non-NASH nonalcoholic fatty liver disease (NAFLD)²⁰⁵
- NAFLD as an aetiology may have lower health-related quality-of-life (HRQOL) scores than other aetiologies of liver disease^{202,203}
- Patients with cirrhosis have poorer scores than patients with no or some fibrosis^{202,203}
- Weight loss in people with NAFLD may improve QOL scores²⁰⁴

Mental health scores

- Not affected by degree of necroinflammatory injury²⁰²
- Not affected by fibrosis stage or parenchymal remodelling (that is, cirrhosis)²⁰²

Bariatric surgery

- Improves overall survival in obesity^{170,204–207}
- Is safe in patients with NASH⁸
- Improves necroinflammatory disease, steatosis and fibrosis¹⁷⁴

Box 4 | Clinical goals to diagnose and manage NAFLD

Diagnostic needs

- Validation of newer non-invasive detection methods to replace the use of invasive, expensive and imperfect 'gold-standard' liver biopsy for the detection and severity evaluation of nonalcoholic steatohepatitis (NASH) and nonalcoholic fatty liver disease (NAFLD), including advanced fibrosis
- Serum-based algorithms as markers for liver injury and fibrosis^{19,229–233}
- Imaging-based system to detect the constellation of lesions that collectively comprise NASH

Management

- Lifestyle intervention, including management of meal content and portions, and exercise programmes
- Therapeutic treatments based on a better understanding of the mechanisms of liver injury from the underlying and closely related links of overweight, obesity and insulin resistance

NASH and HCC

Another consequence of liver disease in patients with NASH is HCC. The incidence of HCC is increasing in both the United States and worldwide. In western countries, approximately 20% of HCCs are found in patients with NASH-related liver disease; and even more if HCCs occurring in the setting of cryptogenic cirrhosis are included, which often represents 'burned out' NASH that has lost its classic histological features (FIG. 1)^{215,216}. A recent study in the United States based on Surveillance, Epidemiology, and End Results (SEER)-Medicare registries from 2004 to 2009 found NASH to be the underlying cause of HCC in 14.1% of nearly 5,000 cases, following alcoholic liver disease in 16% and hepatitis C virus infection in 54.9%. While this represented a 9% annual increase in NASH-related HCC, NASH-HCC cases represented only 5% of liver transplantations for HCC during the same time period, probably reflecting the burden of co-morbid illness in this population. Furthermore, patients with NASH and HCC were older and had shorter survival if they did not receive liver transplantation than other aetiologies of liver disease²¹⁷.

Welzel and colleagues estimated the population attributable fraction (PAF), otherwise stated as the proportion of cases that can be attributed to specific risk factors, in patients with HCC from various causes of liver disease using the SEER-Medicare databases from 1994 to 2007 (REF. 218). Analysis of a cohort of 6,991 people showed that those with obesity and diabetes (presumably with NAFLD) had the highest PAF of HCC (36.6%) compared with other aetiologies, in descending order of magnitude: alcohol intake (23.5%), hepatitis C virus infection (22.4%), hepatitis B virus infection (6.3%) and other (3.2%). An important limitation of this study is that patients did not have a biopsy diagnosis of NAFLD or NASH. The PAF takes into account disease prevalence; thus, if NAFLD were to be eliminated, the effect on HCC would be substantial. However, the risk estimate of HCC in an individual patient with NAFLD is only in the order of 1.5–2.5 compared with 20–25 in a patient with hepatitis C virus infection. Per annum rates of HCC development in patients with NASH are lower than for patients with hepatitis C virus infection (2.6 versus 4)²¹.

As NASH is so prevalent, it is not surprising that HCC as an indication for liver transplantation has increased 3.64-fold in patients with underlying NASH and a BMI of >30 kg per m². In comparison, HCC due to hepatitis C virus has increased only 2.25-fold over a 10-year period from 2002–2012. Whereas diabetes and obesity are independent risk factors for HCC, the increase in NASH-related HCC persisted after controlling for these variables²¹⁹. Although older age, obesity, T2DM and male sex are risk factors for the development of HCC, NASH does seem to contribute independently^{218,220}.

Cirrhosis creates a milieu that is permissive for the development of HCC; thus, current screening recommendations focus primarily on patients with cirrhosis. However, reports of HCC occurring in the absence of cirrhosis are on the rise, with some even developing in the absence of profound fibrosis. By some estimates, 10–75% of cases of HCC may occur in non-cirrhotic NAFLD^{221,222}. This is a serious problem because of the large number of patients affected by NAFLD and NASH at a population level. The individual risk of HCC in patients without cirrhosis who have NAFLD is even less well understood. Consequently, although some individuals at risk for HCC may not be identified at an early stage of malignancy, there is presently insufficient evidence to alter current screening guidelines. Optimal recommendations for HCC screening in patients with NAFLD may change in the future as more is learnt about individual risk in this challenging population.

Outlook

Both the short-term and the long-term outlook for NAFLD will evolve as clinical and basic research continue to re-define the field (BOX 4). NAFLD is an extremely complex and nuanced disease that represents the convergence of many pathways, risk factors and external influences that are not uniform in all patients. Therefore, it seems likely that NAFLD represents a constellation of various phenotypes that will require study from many angles as illustrated in this Primer. NAFLD is highly prevalent, yet only a minor subset of patients progress to advanced liver disease, including HCC. Although many patients with NAFLD have a common metabolic profile (T2DM, hypertension and obesity, among others), not all patients do. The underlying pathophysiology of NAFLD and in particular NASH is strongly linked to insulin resistance, aberrant hepatic lipid metabolism, visceral adiposity and inflammation. However, several other important modulators of disease, such as the environment and diet, can further modify triggers of chronic extrahepatic and intrahepatic immune pathways²²³ and the pathogenetic roles of gut dysbiosis^{224–226}. The extent to which these factors drive disease progression and outcomes will be clarified as we gain further insight into the amassing data in this increasingly important area of research.

Furthermore, differences in genetic susceptibility as well as epigenetics may help to better understand at-risk populations. Therefore, a more in-depth evaluation of these differences will allow us to better understand the phenotypes of NAFLD. In turn, a more sophisticated ability to phenotype patients with NAFLD will further clarify

differences in disease progression between patients and permit a more personalized approach to therapy.

Animal models of disease have offered tremendous insight into relevant mechanisms, but to date, no animal model has faithfully replicated all aspects of the human condition. However, this benchmark may not be compulsory or attainable given the heterogeneity of the human disease.

Just as the pathophysiology of NAFLD is not the same in all patients, treatment will need to be individualized. With rare exception, a commitment to lifestyle modification should be the foundation of any treatment plan, but this is not sufficient for most patients. In the morbidly

obese or in those with severe co-morbid illness, bariatric surgery may have substantial benefit. Many patients will require more-intense intervention — behavioural and pharmacological.

Until recently, the landscape of clinical trials for NASH was somewhat limited. Currently there are hundreds of clinical trials in NASH. The breadth of mechanisms being targeted reflects the complexity of the disease. As the field continues to advance and patients can be better classified by constellations of risk factors, or sub-classifications using genomics or metabolomics, we will be able to offer a more personalized, mechanistically derived treatment approach to each patient.

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Acknowledgements

E.B. is a member of the EPoS (Elucidating Pathways of Steatohepatitis) consortium funded by the Horizon 2020 Framework Programme of the European Union under Grant Agreement 634413.

Author contributions

Introduction (E.M.B.); Epidemiology (V.W.-S.W., V.N. and E.M.B.); Mechanisms/pathophysiology (C.P.D., S.S. and J.J.M.); Diagnosis, screening and prevention (E.B. and C.B.S.); Management (B.A.N.-T.); Quality of life (M.E.R.); Outlook (M.E.R.); Overview of Primer (E.M.B.).

Competing interests

E.M.B. has received consulting fees from Eli Lilly and acted as a paid study pathologist for Rottapharm. C.B.S. has received research grants from GE Healthcare and Siemens. V.W.-S.W. is or has been an advisory board member for Gilead and Janssen, a consultant for AbbVie, Merck and NovoMedica, and has received lecture fees from AbbVie, Echosens and Gilead. B.A.N.-T. has received consulting fees from Nimbus Therapeutics, Bristol-Myers Squibb, Janssen, Conatus, Scholar Rock, Novartis, Galmed, Zafgen and Pfizer. All other authors declare no competing interests.